| | Application No. | Applicant(s) |
|---|---|--|
| | 10/614,954 | HOHN ET AL. |
| Notice of Allowability | Examiner | Art Unit |
| | Medina A. Ibrahim | 1638 |
| The MAILING DATE of this communication appeal claims being allowable, PROSECUTION ON THE MERITS IS herewith (or previously mailed), a Notice of Allowance (PTOL-85) NOTICE OF ALLOWABILITY IS NOT A GRANT OF PATENT RI of the Office or upon petition by the applicant. See 37 CFR 1.313 | (OR REMAINS) CLOSED in this ap or other appropriate communicatio IGHTS. This application is subject | oplication. If not included n will be mailed in due course. THIS |
| 1. \boxtimes This communication is responsive to <u>the amendment of 05</u> | <u>//09/05</u> . | |
| 2. The allowed claim(s) is/are <u>1-5</u> . | | |
| 3. The drawings filed on <u>07 July 2003</u> are accepted by the Ex | aminer. | |
| 4. Acknowledgment is made of a claim for foreign priority ur a) All b) Some* c) None of the: 1. Certified copies of the priority documents have 2. Certified copies of the priority documents have 3. Copies of the certified copies of the priority documents have International Bureau (PCT Rule 17.2(a)). * Certified copies not received: | been received. been received in Application No. | • |
| Applicant has THREE MONTHS FROM THE "MAILING DATE" noted below. Failure to timely comply will result in ABANDONM THIS THREE-MONTH PERIOD IS NOT EXTENDABLE. | | complying with the requirements |
| 5. A SUBSTITUTE OATH OR DECLARATION must be subm INFORMAL PATENT APPLICATION (PTO-152) which give | | |
| 6. CORRECTED DRAWINGS (as "replacement sheets") mus | t be submitted. | |
| (a) ☐ including changes required by the Notice of Draftspers | | -948) attached |
| 1) hereto or 2) to Paper No./Mail Date | | |
| (b) including changes required by the attached Examiner's Paper No./Mail Date | s Amendment / Comment or in the G | Office action of |
| Identifying indicia such as the application number (see 37 CFR 1. each sheet. Replacement sheet(s) should be labeled as such in t | | |
| 7. DEPOSIT OF and/or INFORMATION about the deposit attached Examiner's comment regarding REQUIREMENT | sit of BIOLOGICAL MATERIAL FOR THE DEPOSIT OF BIOLOGIC | must be submitted. Note the CAL MATERIAL. |
| Attachment(s) 1. ☐ Notice of References Cited (PTO-892) | 5. ☐ Notice of Informal F | Patent Application (PTO-152) |
| 2. Notice of Draftperson's Patent Drawing Review (PTO-948) | 6. ⊠ Interview Summary | (PTO-413), |
| 3. ☐ Information Disclosure Statements (PTO-1449 or PTO/SB/0 | Paper No./Mail Da | |
| Paper No./Mail Date 4. Examiner's Comment Regarding Requirement for Deposit | 8. X Examiner's Statem | ent of Reasons for Allowance |
| of Biological Material | 9. Other | |
| | | |
| | | |

Application/Control Number: 10/614,954

Art Unit: 1638

EXAMINER'S AMENDMENT

An examiner's amendment to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

Authorization for this examiner's amendment was given in a telephone interview with Gregory Warren on 06/25/05.

The application has been amended as follows:

In The Title:

The title has been replaced with the following:

--- ISOLATED NUCLEIC ACIDS ENCODING TRICHOTHECENE 3-0-ACETYLTRANSFERASES ----.

In The Abstract:

The abstract has been replaced with the following:

---- The present invention discloses isolated polynucleotides encoding polypeptides having trichothecene 3-0-acetyltransferase activity, recombinant vectors and host cell comprising said polynucleotides----.

In The claims:

In claims 3-5, "gene" has been replaced with --- DNA----.

Reasons for allowance

The following is an examiner's statement of reasons for allowance: the prior art does not teach or reasonably suggest a nucleotide sequence that is 100% identical to

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SEQ ID NO: 5 or a nucleotide sequence encoding SEQ ID NO: 6. The closest prior art is Kimura et al (J. Biol. Chem. (1998), 273 (3): 1654-1661, Applicant's IDS) who teach a nucleotide sequence having 99% sequence identity with Applicant's SEQ ID NO: 5. See attached sequence alignment results (Exhibit A/B) between SEQ ID NO: 5/6 and the prior art sequences.

Any comments considered necessary by applicant must be submitted no later than the payment of the issue fee and, to avoid processing delays, should preferably accompany the issue fee. Such submissions should be clearly labeled "Comments on Statement of Reasons for Allowance."

Contact information

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Medina A. Ibrahim whose telephone number is (571) 272-0797. The Examiner can normally be reached Monday -Thursday from 8:00AM to 5:30PM and every other Friday from 9:00AM to 5:00 PM. Before and after final responses should be directed to fax nos. (703) 872-9306 and (703) 872-9307, respectively.

If attempts to reach the Examiner by telephone are unsuccessful, the Examiner's supervisor, Dr. Amy Nelson, can be reached at (571) 272-0804.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

For all other customer support, please call the USPTO Call Center (UCC) at 800-786-9199.

6/25/05 Mai

MEDINA A. IBRAHMA
PATENT EXAMINER
PATENT EXAMINER
1638

06 May 2005

Alignment Results

Alignment: Global DNA alignment against reference molecule Parameters: Scoring matrix: Linear (Mismatch 2, OpenGap 4, ExtGap 1)

Reference molecule: SEQIDNO:5, Region 1-1356
Number of sequences to align: 2
Settings: Similarity significance value cutoff: >= 60%



| Summary of Percent Matches: | | | | | |
|-----------------------------|-------|------|----|-----------|-----|
| Reference: SEQIDNO:5 | 1 - | 1356 | (| 1356 bps) | |
| Keleleuce: Protogram | 126 | 1490 | i | 1356 bps) | 99% |
| Sequence 2: Kimura Tril01 | 135 - | 1420 | ١, | Taan pbal | 220 |

| SEQIDNO:5 | 1 atggettteaagatacagetegacaceeteggeeagetaceaggeeteetttegatetac |
|---------------------------|--|
| Kimura Tril0 | 135 atggettteaagatacagetegacaceeteggeeagetaceaggeeteetttegatetac |
| SEQIDNO:5 Kimura Tri10 | 61 acccaaatcagtctcctctaccccgtctctgattcctctcaatatcccactattgtcagc 195 acccaaatcagtctcctctaccccgtctctgatccctctcaatatcccactattgtcagc |
| SEQIDNO:5 | 121 accttcgagcaaggtcttaagcgcttctccgaagccgtcccatgggtcgcaggccaggtc |
| Kimura Tri10 | 255 accttcgagcaaggtcttaagcgcttctccgaagccgtcccatgggtcgcaggccaggtc |
| SEQIDNO:5 | 181 aaagccgagggcattagcgagggaaacacaggaacttcctttatcgtcccttttgaggac |
| Kimura Tri10 | 315 aaagccgagggcattagcgagggaaacacgggaacttcctttatcgtcccttttgaggac |
| SEQIDNO:5 | 241 gttcctcgtgttgtagtgaaagacctccgcgatgatccttcagcgcccacgatcgagggt |
| Kimura Tri10 | 375 gttcctcgtgttgtagtgaaagacctccgcgatgatccttcagcgcccacgatcgagggt |
| SEQIDNO:5 | 301 atgagaaaggcgggataccctatggcgatgtttgacgagaacatcatcgcgccaaggaag |
| Kimura Tri10 | 435 atgagaaaggcgggataccctatggcgatgtttgacgagaacatcatcgcgccaaggaag |
| SEQIDNO:5 | 361 acgttacetattggacetggtactggtcccgacgacccaaagcctgtaattctattgcag |
| Kimura Tri10 | 495 acgttacetattggacetggtactggccccgacgacccaaagcctgtgattctattgcag |
| SEQIDNO:5 | 421 ctcaacttcatcaagggcggactcatcctcactgtcaacggacagcacggtgctatggat |
| Kimura Tri10 | 555 ctcaacttcatcaagggcggactcatcctcactgtcaacggacagcacggtgctatggat |
| SEQIDNO:5 Kimura Tri10 | 481 atggtaggccaagatgcggtgatccgtctactctccaaggcgtgccgtaacgacccattc 615 atggtaggccaagatgcggtgatccgtctactctccaaggcgtgccgtaacgacccattc |
| SEQIDNO:5 Kimura Tri10 | 541 accgaagaggaaatgacggccatgaacctcgatcgcaagacgatagttccttaccttgaa 675 accgaagaggaaatgacggccatgaacctcgatcgcaagacgatagttccttaccttgaa |
| SEQIDNO:5 Kimura Tril0 | 601 aactatacgattggccccgaggtagatcatcagattgtcaaagctgatgtagctggtg735 aactacacgattggccccgaggtagatcatcagattgtcaaacctgatgtagctggtggt |
| SEQIDNO:5 | 661 gacgetgtteteacgeeggteagtgeaagetgggegttetteacatteageeceaaggee |
| Kimura Tri10 | 795 gacgetgtteteacgeeggteagtgeaagetgggegttetteaaatteageeceaaggee |
| SEQIDNO:5 | 721 atgtcagagetcaaggatgetgetaccaagactettgacgcatcaacaaagttegtgteg |
| Kimura Tri10 | 855 atgtcagagetcaaggatgetgetaccaagactettgacgcatcaacaaagttegtgteg |
| SEQIDNO:5 Kimura Tril0 | 781 actgacgatgctctttcggcgttcatctggaaatcggcctctcgcgtgcgt |
| SEQIDNO:5 | 841 atcgatggctctgcacctaccgagttctgccgtgctgttgatgctcgaccggcaatgggt |
| Kimura Tri10 | 975 atcgatggctctgcacctaccgagttctgccgtgctgttgatgctcgaccggcaatgggt |
| SEQIDNO:5 Kimura Tri10 | 901 gtctcgaacaactacccaggccttcttcaaaacatgacctaccacaactcgaccatcggc 1035 gtctcgaacaactacccaggccttcttcaaaacatgacctaccacaactcgaccatcggc |
| SEQIDNO:5 | 961 gaaatcgccaacgagtcactcggcgcaacagcatcacgccttcgttcagaactcgaccc |
| Kimura Tri10 | 1095 gaaatcgccaacgagtcactcgqcgcaacagcatcacgccttcgttcagaactcgaccc |

| SEQIDNO:5 | 1021 gcgagcatgcgccagcgaacaagaggtctcgcgacgtacctgcacaacaaccccgacaag |
|---------------------------|---|
| Kimura Tri10 | 1155 gcgagcatgcgccagcgaacaagaggtctcgcgacgtacctgcacaacaaccccgacaag |
| SEQIDNO:5 Kimura Tri10 | 1081 tccaacgtatccctgacggctgatgcggacccatctaccagcgtcatgctgagttcttgg 1215 tccaacgtatctctgacggctgatgcggacccatctaccagcgtcatgctgagttcttgg |
| SEQIDNO:5 | 1141 gccaaggtgggactctgggattacgactttgggctcggactgggtaagcccgagactgtg |
| Kimura Tri10 | 1275 gccaaggtcggactctgggattacgactttgggttcggactgggtaagcccgagactgtg |
| SEQIDNO:5 | 1201 agacggccaatctttgagcctgttgagagcttgatgtactttatgcccaagaagcctgat |
| Kimura Tri10 | 1335 agacggccaatctttgagcctgttgagagcttgatgtactttatgcccaagaagcctgat |
| SEQIDNO:5 | 1261 ggcgagttctgtgcggcgctttctctgagggatgaggatatggaccgattgaaggcggat |
| Kimura Tri10 | 1395 ggcgagttctgtgcggcgctttctctgagggatgaggatatggaccgattgaaggcggat |
| SEQIDNO:5 | 1321 aaggagtggaccaagtatgcgcagtacgttggttag |
| Kimura Tri10 | 1455 aaggagtggaccaagtatgcgcagtacgttggttag |

Exhibit B

99%

06 May 2005

Alignment Results

Global Protein alignment against reference molecule Alignment:

Parameters: Scoring matrix: BLOSUM 62

Reference molecule: SEQIDNO:6, Region 1-451

Number of sequences to align: 2

Reference: SEQIDNO:6
Sequence 2: Kimura Tri101 Pro SEQIDNO: 6

Settings: Similarity significance value cutoff: >= 60%

Summary of Percent Matches:

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451



